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## IN THE CLATMS

Please amend claims 1-5 and add claims 35-36 as follows:

- (amended) A method for creating target single strand regions in a plurality of double stranded DNA molecules for use in joining the DNA molecules, comprising:
- (a) nicking at least two sites bordering a target region within the DNA molecules with at least one site-specific nicking endonuclease;
- (b) subjecting the nicked DNA molecules from step (a) to conditions that selectively denature the target region to create the target single stranded region; and
- (c) joining the DNA molecules from step (b) by means of the target single strand regions.
- 2. (amended) A method of claim 1 wherein the at least two sites bordering the target region are located on a single strand of the double stranded DNA so that the target single stranded region comprises a gap in the double stranded DNA.
- 3. (Ameded) A method for creating a target single strand region at a terminus of a linear double stranded DNA molecule for use in joining the DNA molecule to a second DNA molecule by means of the single strand region, or for detecting, purifying or selectively mutagenizing the DNA molecule, comprising

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- (a) nicking at least one site bordering the target region at the terminus of the linear double stranded DNA with at least one sitespecific nicking endonuclease;
- (b) subjecting the nicked DNA molecules from step (a) to conditions that selectively denature the target region to create the target single stranded region; and
- (c) joining the DNA molecule to a second DNA molecule by means of the single strand region, or detecting, purifying or selectively mutagenizing the DNA molecule by means of the single strand region.
- 4. (Amended) The method of claim 3 wherein the DNA terminus is pre-existing.
- 5. (Amended) The method of claim 3 wherein the DNA terminus is formed by site-specific endonuclease cleavage.
- 35. (New) A method for creating target single strand regions in a double stranded DNA molecule for use in detecting, purifying or selectively mutagenizing the DNA molecule, the method comprising:
- (a) nicking at least two sites bordering a target region in the DNA molecule with at least one site-specific nicking endonuclease;
  and
- (b) subjecting the nicked DNA molecules from step (a) to conditions that selectively denature the target region for creating the target single stranded region; and